

- a a region that doesn't interact with clone 10 (NMP1), 9 and 15 in the 2-hybrid test
- b a region that interact with these clones as strongly as the full-size TRAF2
 - - RING finger motif
 - ☐ TFIIIA-like zinc-finger motifs
 - 置 N-TRAF
 - E C-TRAF

FIG. 1

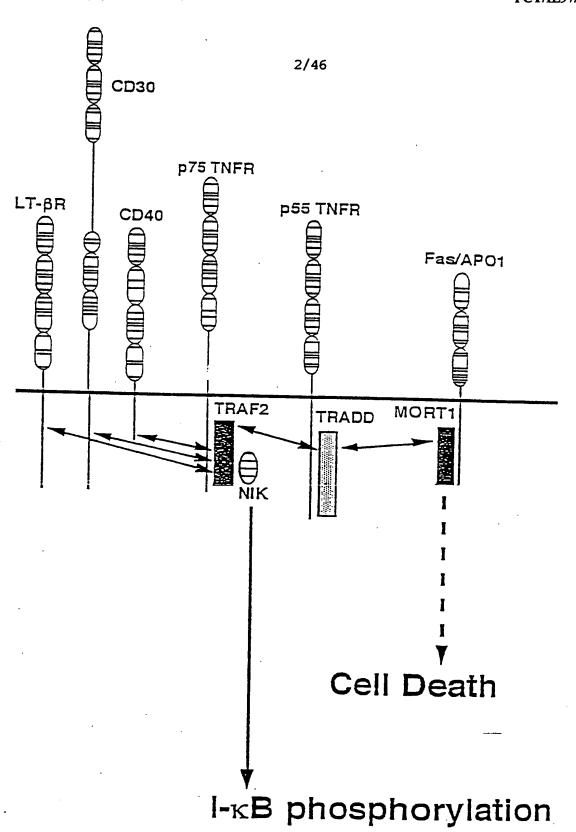


FIG. 2A



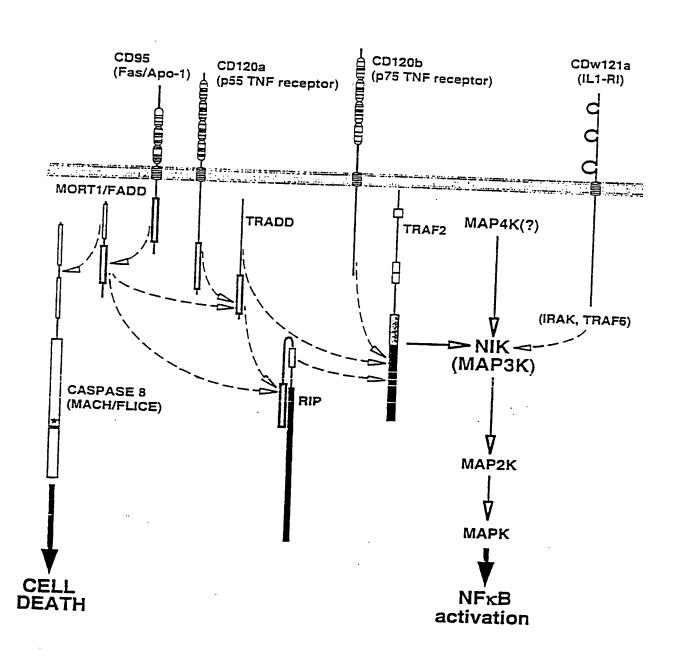


FIG. 2B

09/155676 PCT/IL97/00117

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Le	ngth: 1906	July 7, 1	996 12:35	Type: N Ch	eck: 7122 .
1	CATTGGGTCA	CGCGGTGGCG	GCGCTCTAGA	ATAGTGGATC	CCCCGGGCTG
51	CAGGAATTCG	ATTCGAGGCC	ACGAAGGCCG	GCGGCGCGC	GCAnGCACCG
101	GCCCGGGGAn	AGGCnCCATG	AGCGGATCnC	nGAACnATGA	CAAAAGACAA
151	TTTCTGCTGG	AGCGACTGCT	GGATGCAGTG	AAACAGTGCC	AGATCCGCTT
201	TnGAGGGAGA	AAGGAGATTG	CCTCGGATTC	CGACAGCAGG	GTCACCTGTC
251	TGTGTGCCCA	GTTTGAAGCC	GTCCTGCAGC	ATGGCTTGAA	GAGGAGTCGA
301	GGATTGGCAC	TCACAGCGGC	AGCGATCAAG	CAGGCAGCGG	GCTTTGCCAG
351	CAAAACCGAA	ACAGAGCCCG	TGTTCTGGTA	CTACGTGAAG	GAGGTCCTCA
401	ACAAGCACGA	GCTGCAGCGC	TTCTACTCCC	TGCGCCACAT	CGCCTCAGAc
451	gTGGGCCGGG	GTCGCGCCTG	GCTGCGCTGT	GCCCTCAACG	AACACTCCCT
501	GGAGCGCTAC	CTGCACATGC	TCCTGGCCGA	CCGCTGCAGG	CTGAGCACTT
551	TTTATGAAGA	CTGGTCTTTT	GTGATGGaTG	AAGAAAGGTC	CAGTATGCTT
601	CCTACCATGG	CAGCAGGTCT	GAACTCCATA	CTCTTTGCGA	TTAACATCGA
651	CAACAAGGAT	TTGAACGGGC	AGAGTAAGTT	TGCTCCCACC	GTTTCAGACC
701	TCTTAAAGGA	GTCAACGCAG	AACGTGACCT	CCTTGCTGAA	GGAGTCCACG
751	CAAGGAGEGA	GCAGCCTGTT	CAGGGAGATC	ACAGCCTCCT	CTGCCGTCTC
801	CATCCTCATC	AAACCTGAAC	AGGAGACCGA	CCCTTGCCTG	TCGTGTCCAG
851	GAATGTCAGT	GCTGATGCCA	AATGCAAAAA	GGAGCGGAAG	AAGAAAAAGA

FIG. 3A

901	330000000				
					GCAGAACTCT
951 	GGGGACGTGT	TTAAAAAGAC	ACCTGGGGCA	. GGGGAGAGCT	CAGAGGACAA
1001	CTCCGACCGC	TCCTCTGTCA	ATATCATGTC	CGCCTTTCAA	AGCCCCTTCG
1051	GGCCTAACTC	CAATGGAATG	363663663	CCCITIGAL.	ATTGATTCCC
 1101	TCTCTTTT		AGAGCAGCAA	CTCATGGAAA	ATTGATTCCC
	IGICITIGAA	CGGGGAGTTT	GGGTACCAGA	AGCTTGATGT	GAAAAGCATC
1151	GALGALGAAG	ATGTGGATGA	AAACGAAGAT	GACGTGTATG	GAAACTCATC
1201	AGGAAGGAAG	CACAGGGGCC	ACTCGGAGTC	GCCCGAGAAG	CCACTGGAAG
1251	GGAACACCTg	CCTCTCCcAG	ATGCACAGCT	GGGC+CCCC	Care -
1301	CaCAaTGACT	CCG3C3TCCT	CERTAGO		GAAGGTGCTG
1351	CCC2 CC2	CCGACATCCT	CITCCCTGTC	AGTGGCGTGG	gCTCCTACAG
		gCCCCCTCG	GAAGCCTGGA	GAACGGGACA	GGACCAGAGG
1401	ACCACGTTCT	CCCGGATCCT	GGACTTCGGT	ACAGTGTGGA	AGCCAGCTCT

FIG. 3B

1901 AAAAAA

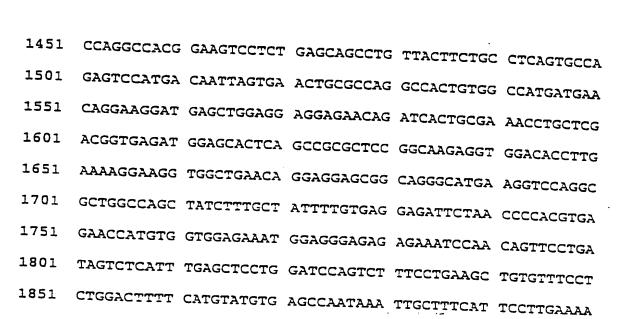


FIG. 3C

TRANSLATE of: 9hhh check: 7122 from: 1 to: 1906 generated symbols 1 to: 635.

9hhh.pep Length: 604 August 23, 1996 15:03 Type: P Check: 4554 1 XTGPGXGXMS GSXNXDKRQF LLERLLDAVK QCQIRFXGRK EIASDSDSRV 51 TCLCAQFEAV LQHGLKRSRG LALTAAAIKQ AAGFASKTET EPVFWYYVKE 101 VLNKHELQRF YSLRHIASDV GRGRAWLRCA LNEHSLERYL HMLLADRCRL 151 STFYEDWSFV MDEERSSMLP TMAAGLNSIL FAINIDNKDL NGQSKFAPTV 201 SDLLKESTON VTSLLKESTO GVSSLFREIT ASSAVSILIK PEQETDPCLS 251 CPGMSVLMPN AKRSGRRKRK *PT*SHLMMR KMSRTLGTCL KRHLGQGRAQ 301 RTTPTAPLSI SCPPLKAPSG LTPMESEQQL MEN*FPVFER GVWVPEA*CE KHR**RCG*K RR*RVWKLIR KEAQGPLGVA REATGREHLP LPDAQLGSAE 351 401 GAAQ*LRHPL PCQWRGLLQP SRCPPRKFGZ RDRTRGPRSP GSWTSVQCGS QLSRPRKSSE QPVTSASVPE SMTISELRQA TVAMMNRKDE LEEENRSLRN 501 LLDGEMEHSA ALRQEVDTLK RKVAEQEERQ GMKVQALASY LCYFVRRF*P 551 HVRTMWWRNG GREKSNSS** SHLSSWIQSF LKLCFLWTFH VCEPINCFHS 601 LKKK

clone 10	Length: 26	31 August	23, 1996 17	7:18 Type:	N Check: 5107
1	CCCCTCTCAC	AGCCCAGGCC	ATCCAAGAGG	GGCTGAGGA	A AGAGCCCATC
51	CACCGCGTGT	cTGCAGcGGA	GCTGGGAGG	AAGGTGAAC	C GGGCACTACA
101	GCAAGTGGGA	GGTCTGAAGA	GCCCTTGGAC	GGGAGAATA:	r aaagaaccaa
151	GACATCCACc	GCCAAATCAA	GCCaAtTACC	ACCAGACCC	CCATGCCCAg
201	CCGAGAGAGC	TETCGCCAAG	GGCCCcAGGG	CCCCGGCCA	CTGAGGAGAC
251	AACAggCAGA	GCCCCTAAGC	TCCAGCcTCC	Teteceace:	A GAGCCCCAG
301	AGCCAAACAA	GTCTCCCC	ttGACTttGA	CCAAGGAGGA	GTCTGGGATG
351	TGGGAACCCT	TACCTCLGTC	CTCCCTGGAG	CCAGCCCCTC	CCAGAAACCC
401	CAGCTCACCA	GAGCGGAAAG	CAACCGTCCC	GGAGCAGGA	CTGCAGCAGC
451	TGGAAATAGA	ATTATTCCTC	AACAGCCTGT	CCCAGCCATI	TTCTCTGGAG
501	GAGCAGGAGC	AAATTCTcTc	GTGCCTCAGC	ATCGACAGCC	TCTCCCtGTC
551	GGATGACAGT	GAGAAGAACC	CATCAAAGGC	CTCTCAAAGC	TCGCGGGACA
601	CCCTGAGCTC	AGGCGTACAC	TCCTGGAGCA	. GcCAGGCCGA	GGcTCGAAGc
651	TCCAGCTGGA	ACATGGTGcT	GGCCCGGGgg	CGGCCCACCG	ACACCCAAG
701	CTATTTCAAT	GGTGTGAAAG	TCCAAATACA	GTCTCTTAAT	GGTGAACACC
751	TGCACATCCG	GGAGTTCCAC	CGGGTCAAAG	TGGGAGACAT	CGCCACTGGC
801	ATCAGCAGCC	AGATCCCAGC	TGCAGCCTTC	AGCTTGGTCA	CCAAAGACGG
851	GCAGCCTGTT	CGCTACGACA	TGGAGGTGCC	AGACTCGGGC	ATCGACCTGC
901	AGTGCACACT	GGCCCCTGAT	GGCAGCTTCG	CCTGGAGCTG	GAGGGTCAAG
951	CATGGCCAGC	TGGAGAACAG	GCCCTAACCC	TGCCCTCCAC	CGCCGGeTCC
1001	ACACTGCCGG	aAAGCAGCCT	TCCTGCTCGG	tGCACGATGC	TGCCCTGaAA
1051	Acacaggete	AGCCGTTCCC	Aggggatytg	.CCAGcCCCC	cGGcTcArcA
1101	G.tGGGaAcC	AGGGCCTCG.	CAGC.AGC.A	AGGT.gGGGG	CAAGC.AGAA
1151					CTGCTGaadA
1201	AAAcayt.cc	GCcAcGtGAA	Gagacagaag	GAGGATGG.C	AGGAgTtA
1251	CcTygGGGAA	aCaAAAcAGg	gaTcTTt.tT	cTGcCCcTGc	TCCAGT.cGA

FIG. 4A

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1301	gtTGGCCTG. ACCCGCTTGG A.TCAgtGAC CATTTGtTGG CAGA.CAGGG
1351	GagAgCAGCT TCCAGCCTGG gTCAGAAGGG GTGGGCGAGC CCtTcGGCCC
1401	CTCACCCT.c CAGGCTGCTG tG.AGAGTGT CAAGTGtGTA AGGG.CCCAA
1451	A. CTCAGG. T TCAGTGCAGA ACCAGGT.CA GCAGGTATGC CCGCCCG.TA
1501	GGTTAAGG GGGCCETET. AAACCCCTTG eCT.GGCCT. CAECT.GGCC
1551	AGCTCA.CCC CTTTTGGGTG TAGGGGAAAA GAATGCCTGA CCCTGGGAAG
1601	GCTWCCCTGG TAGAATACAC CACACTTTTC AGGTTGTTGC AACACAGGTC
1651	CTGAGTTGAC CTCTGGTTCA GCCAAGGACC AAAGAAGGTG TGTAAGTGAA
1701	GTGGTTCTCA gT.CCCCAGA CATGTGCCCC TTTGCTGCTG GCTACCACTC
1751	TTCCCCAGAG CAGCAGGCCC CGAGCCCCTT CAGGCCCAGC ACTGCCCCAG
1801	ACTCGCTGGC ACTCAGTTCC CTCATCTGTA AAGGTGAAGG GTGATGCAGG
1851	ATATGCCTGA CAGGAACAGT CTGTGGAŁGG ACATGATCAG TGCT.AAGG.
1901	AAAGCAGCAG AGAGAGACGY TCCGGCGCCC CAg.CCCCAC T.ATCAGTgT
1951	.CCAGCGTGC T.GGTT.CCC CAG.AGCACA GCT.CAG.CA TCA.CACTGA
2001	CACT.CACCC T.GCCCTGCC CCT.GGCCA. GAGGGTACTG CCG.ACGGCA
2051	CTTTGCAc.T CTGATG.ACC TCAAAGCACT TTCATGgcT. GcCCTctG
2101	GCAGGG.CAG GG.CAGGG.C AGTGACA.CT GTAGG.AGCA TA.gCAA.GC
2151	CAGGAGATGG GGTG.AAGGG A.CACAGTCT TGAGCTGTCC A.CATGCATG
2201	TGACT.CCTC AAACCTCTTCCAG.ATTT CTCTAAGAAT AGCA.CCCCC
2251	TT.CCCCATT-GCCCCAGCTT AGCCTCTTCT CCCAGGGGAG CTA.CTCAGG
2301	ACTCACGTAG CATTAAATCA GCTGTG.AAT CGTCAGGGGG TGTCTGCTAG
2351	CCTCAACCTC CTGGGGCAGG GGACGCCGAG ACTCCGTGGG AGAAGCTCAT
2401	TCCCACATCT TGCCAAGACA GCCTTT.GTC CAGCTGTCCA CATTGAGTCA
2451	gACTGCTCCC GGGGAGAGAG CCCCGGCCCC CAGCACATAA AGAACTGCAG
2501	CCTTGGTACT GCAGAGTCTG GGTTGTAGAG AACTCTTTGT AAGCAATAAA
2551	GTTTGGGGTG ATGACAAATG TTAAAAAAAG GCCTTCGTGG CCTCGAATCA
2601	AGCTTATCGA TACCGTCGAC CTCGAGGGGG G

FIG. 4B



L	ength: 1253	July 10, 1	996 clonel	5	
1	CATTGGAGTC	ACGCGGTGGC	GGCGCTCTAG	AATAGTGGAT	CCCCGGGCTg
51	CA.GGAATTC	GATTCGAGCC	CACGAAGGCC	CCTTCTTCTG	TGGTCGCGGC
101	ACGTTTACaG	CCGCAAGCAc	CCAGCGGCAg	CTGAAGGAGG	CTTTTGAgAG
151	GCTCCTGCCC	CAGGTGGAGG	CGGCCCGCAA	GGCCATCCgC	GCCGCTCAGG
201	TGGAGCGCTA	TGTGCCCGAA	CACGAGCGAT	GCTGCTGGTG	CCTGTGCTGC
251	GGCTGTGAGG	TGCGGGAACA	CCTGAGCCAT	GGAAACCTGA	CGGTGCTGTA
301	CGGGGGGCTG	CTGGAGCATC	TGGCCAGCCC	AGAGCACAAG	AAAGCAACCA
351	ACAAATTCTG	GTGGGAGAAC	AAAGCTGAGG	TCCAGATGAA	AGAGAAGTTT
401	CTGGTCACTC	CCCAGGATTA	TGCGCGATTC	AAGAAATCCA	TGGTGAAAGG
451	TTTGGATTCC	TATGAAGAAA	AGGAGGATAA	AGTGAtCAAG	GAGaTGgCAG
501	CTCAGATCCG	TGaGGTGGAg	CAGAGCCGAC	AGGAGGTGGE	TCGGtCTGTc
551	TTAGAGCCTC	AGGCAGTGcC	AGACCCAGAA	GAGGGCTCTT	CAGCACCTAG
601	AAGCTGGAAA	GGGATGAACA	GCCAAGTAGC	TTCCAGCTTA	CAGCAGCCCT
651		CCTGCCACCA	GCTCCAGAGC	TTGAcTGGAT	GGAGACAGGA
701	CCATCTCTGA	CATTCATTGG	CCATCAGGAT	ATACCAGGAG	TTGGTAACAT
751	CCACTCAGGT	GCCACACCTC	CCTGGATGAT	CCAAGATGAA	GAATACATTG
801	CTGGGAACCA	AGAAATAGGA	CCATCCTATG	AAGAATTTCT	TAAAGAAAAG
851	GAAAAACAGA	AGTTGAAAAA	ACTCCCCCA	GACCGAGTTG	GGGCCAACTT
901				GCTGCCCTCT	-
	GTCTGGAATA	, .	•	•	
1001	•		•	CAGAAAAAAG	•
1051	•••		•	AAAGTCAACC	•
1101	• • •			TCTTTCTTAG	•
	TACTCATTCA				•
	CGAATCAAGC	TTATCGATAC	CGtCGACCTC	GAGGGGGGC	CGTACCCACT
1251	TTT				

FIG. 5A

PCT/IL97/00117

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TRANSLATE of: 15cc check: 9389 from: 2 to: 1253 generated symbols 1 to: 417.

401 ESSLSIPSTS RGGRTHF

15cc.pep Length: 417 August 23, 1996 14:32 Type: P Check: 7921 ... 1 IGVTRWRRSR IVDPRAAXNS IRAHEGPFFC GRGTFTAAST QRQLKEAFER 51 LLPQVEAARK AIRAAQVERY VPEHERCCWC LCCGCEVREH LSHGNLTVLY 101 GGLLEHLASP EHKKATNKFW WENKAEVOMK EKFLVTPQDY ARFKKSMVKG 151 LDSYEEKEDK VIKEMAAQIR EVEQSRQEVV RSVLEPQAVP DPEEGSSAPR 201 SWKGMNSQVA SSLQQPSNLD LPPAPELDWM ETGPSLTFIG HQDIPGVGNI 251 HSGATPFWMI QDEEYIAGNQ EIGPSYEEFL KEKEKQKLKK LPPDRVGANF 301 DHSSRTSAGW LPSFGPRLE* WTPLAVQTST PKLKLQQ*RS SHIQKKANHA

351 LYQLP*G*KP KSTKPLLYLP PKFFIIVFLR KQTYSFI*FN KVLFFGLRGL

FIG. 5B

(°.)

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TTG CTA 1 AGC GGG GGG ACT GTG CCG TGT GGA ACG TGT AGC TGT TGA AGG TGG ACT CTG TTA CCA GCA CAC ATA AAC AGG CAG AGA AAG K AGG AAT N CAA CTG TGT TCT GCA TGA 91 GAG 151 GAC 211 GGA TGA GTA TGT GTG GCA TIT CIC CCC CAA CCC AAG GCT TGG TCT CTC GTT TTT GGA GGA CTC TGT ပ္ပ CIG ATG ATG GAC 181 AGT

FIG. 6A



CCA P CGA R CAC H CTG L TTC F AAA K ဥဌဌ GAG E AGC S TAT Y CCG P CCT P CCA P AAC N ည္သည္ ဗ AAA K ACG T TGT C CAG Q AAG K AGC ATC I CAC H CTC L TGT

FIG. 6B

1121 462 TCC AGA TCC GGG GAG CCC AAA ACT GAG GAC GAG GGT CTC CTC ACT GGG T R											ฮ	A																					
TCC AGA TCC CGG GAG CCC AAA ACT GAG GAC AAC GAG GGT CTC CTC AAA CTC AGA CCA GAG CCC AAA ACT GAG GAC CTC TGG CCC CTC AAA CTC AAG CCA GTG GAT TAT GAG TAC CGA GAA GTC CAC TGG CCC ACG CAC R											ŭ	~		٠.			70			75			"			٤.							
TCC AGA TCC CGG GAG CCC AAA ACT GAG GAC AAC GAG GAC CTC AAA CTC AGA CCC AAA ACT GAG GAC CAC TGG CCC CTC AAA CTC AAG CCA GTG GAT TAT GAG TAC CGA GAA GAG GTC CAC TGG CCC ACG A C L C R P V D Y E Y H E E B V H W A T H CGC CTG GGC AGA GCC TCC TTC GGA GGG GTG CAC TGG CCA ACG A G L C A V K K V R L E V F R A E E L CAG TGC CTG GTC TTC GGA GTG CAC GAG ATG GAG GAC GAG CTG CAG TGC CTG GAC AGA ATT GTC CTT TAT CGG CCA GAG GAG GTG A G L T S P R I V P L Y R A E E L CGC CTG GGA TTG CCT TTC ATG GAG TTT CGG CCA GAG GAG GTG GCA GGA TTG ATG GAG TTC TTC TAT GAG TTT CGG CCA GAG GAG GTG A G L T S P R I V P L G A V K R GGC TGT CTC ATG GAG GTG CTG TAC TAT GAA GCT GTG AAG I TAT I F H E L L E G C TAC TAC TAC TAC TAC TAC TAC TAC TAC T		ACT	E		CAG	œ		SGC	t)		ATC	Σ		ິບ	ച											ပ္ပ	o.					-	-
TCC AGA TCC CGG GAG CCC AAA AT GAG GAC AAC GAG GGT CTG 9 R R S R E P S P K 1411 AAA CTC AAG CCA GTG GAT TAT GAG TAC CGA GAA GAA GTC CAC TGG GCC ACG K L L R P R E P V H W A T T L L L R P V D Y E Y R R E P V H W A T T L L G R G S C TCC TTC GAA GTG GTG CAC AGG ATG GAG GAC AAG GAG GAG GAG GAG GAG GAG GA											CIG	1		ეეე	ပ		AAG	×		GGT	ပ		CTC	ם		C.P.	ø	,				AGC	ຮ
TCC AGA TCC CGG GAG CCC AGA ACT GAG GAC AAC GAG GGT GTG S R S R E P S P K T 1411 AAA CTC AGA CCA GTG GAT TAT GAG TAC CGA GAA GAA GTC CAC TGG GCC K L K P V D Y E Y R E E V H W A A CGC CTG GGC AGA GTG CAC TGG GCG GTG GTG CAC AGA GTG CAC AGA GTG GAC AGA GTG CAC TGG GAC AGA GTG GAC GAC GAC GAC GAC GAC GAC GAC GAC GA					ACG (_					GAG	ចា		GAA	വ		GTC	>		GAG	ш		GTG	>		CTT	.ı		ACC	۳		AGC	တ
TCC AGA TCC CGG GAG CCC AAA ACT GAG GAC AAC GAG GGT S R S R E P S P K T 111 AAA CTC AGA CCA GTG GAT TAT GAG TAC CGA GAA GAA GAA GAC GAG AAA CTC AAG CCA GTG GAT TAT GAG TAC CGA GAA GAA GAC GAC GAC CGC CTG GGC AGA GGC TCC TTC GGA GAG GAA GAA GAC GAG CGC CTG GGC AGA GGC TCC TTC GGA GAG GAA GAA GAC GAG CGC CTG GGC AGA GGC TCC TTC GGA GAG GAA GAA GAC GAG A G L T S P R I V P L S R A CAG TGC CCT GTC AAA AAG GTG OGC CTG GAA GTA TTT CGG GCA A G L T S P R I V P L Y G A A A C L T S P R I V P L G G G C CTG GAG GCC A V K K V R C CT C TCG CCC GAG GCC CTG GAC GCC A C C A V K K V R L B L C G G G G C CTG GAC GCC A C C A V K K V R L B L C G G G G C CTG GAC GCC A C C A V K K V R L B L C G G G G G C CTG GAC GCC A C C A V K K V R L B C C G G G G G G G G G G G G G G G G G											GAG	មា		AGA	œ.		CTG	J		ប	J		AAC	z		TGT	ပ		GAG	ш			
TCC AGA TCC GGG GAG CCC AAA ACT GAG GAC AAC GAG S R S R S R T T E D N E AAA CTC AGG CAG GAG CCC AAA ACT GAG GAC AAC GAG S A C C A C C C A C C C C A C C C C C C								GAC	_					GTG	>	•	CAG	ď		ပ္ပ	æ		GAC	Ω		GTG	>		AC.A	۴			
TCC AGA TCC CGG GAG CCC AGC CCC AAA ACT GAG GAC AAC S R		SAG	f.3		.AC	_		AG		•	9	~		CCT	Æ		ပ္ပပ္ပ	ပ		CAG	ø		GCT	æ		GCT	æ						
TCC AGA TCC CGG GAG CCC AGC CCC AAA S R S R E P S P K AAA CTC AAG CCA GTG GAT TAT GAG TAC K L K P V D Y E Y CGC CTG GGC AGA GGC TCC TTC GGA GAG R L G R G S F G E CAG TGC CTG TCC TCC TTC GGA GAG Q C A V K K V R L E E GTC AAC TTC ATG GAG CTG CTG GA V N I F M E L L E GGC TGT CTC CCA GAG GAC CG GC CT G C L P E D R A L TAC CTC CAC TCA CGA AGG TTC TG CAC Y L H S R R I L H AGC GAT GGG AGC CAC GCA GCC TC G C L P E D R D C TAC CTC CAC TCA CGA AGG TTC TG CAC G C L P E D R L C G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC CAC CAC GCA AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC CAC AGG ATT CTG CAC TAC CTC CAC CAC CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC CAC CAC AGG ATT CTG CAC TAC CAC TCAC TCAC CAC CAC CAC CAC CAC CTC TG TAC CAC CAC CAC CAC CAC CAC CAC CAC CAC		AAC	z		ST S	>		ATG	Σ		_			GGA	ဗ		CIG	J		ပ္ပဋ္ဌ	ပ		Æ	×		CAT	×		CCT	Δ,			
TCC AGA TCC CGG GAG CCC AGC CCC AAA S R S R E P S P K AAA CTC AAG CCA GTG GAT TAT GAG TAC K L K P V D Y E Y CGC CTG GGC AGA GGC TCC TTC GGA GAG R L G R G S F G E CAG TGC CTG TCC TCC TTC GGA GAG Q C A V K K V R L E E GTC AAC TTC ATG GAG CTG CTG GA V N I F M E L L E GGC TGT CTC CCA GAG GAC CG GC CT G C L P E D R A L TAC CTC CAC TCA CGA AGG TTC TG CAC Y L H S R R I L H AGC GAT GGG AGC CAC GCA GCC TC G C L P E D R D C TAC CTC CAC TCA CGA AGG TTC TG CAC G C L P E D R L C G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC TCA CGA AGG ATT CTG CAC G C L P E D R A L TAC CTC CAC CAC CAC GCA AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC CAC AGG ATT CTG CAC TAC CTC CAC CAC CAC CAC AGG ATT CTG CAC TAC CTC CAC TCA CAC CAC CAC CAC AGG ATT CTG CAC TAC CAC TCAC TCAC CAC CAC CAC CAC CAC CTC TG TAC CAC CAC CAC CAC CAC CAC CAC CAC CAC		GAC	Ω		GAA	ы		AGG	œ					TAT	> -		TC	လ		CTG			ST C	>		ပ္ပ	೮		ATC	н		AAG	×
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FIG. 6C

AGC CAC H H GCT A A A CGCT E E GCA GGCA A A A A A S S S CTC L ACC AAC AAG K TAC Y CCC ¥ × CTG L AGC S AAT N CCC P GAA E CGG R AGC S GAC D CCC P GAG E TGG W GAG E AAC N ATC I CA O ე ე CCA P ATG M CCA P CTC AGC S CC P P CCA P AAT N TTC F 000 0 TCA S CCC A CTC L TCT S AGC S AGG R GAG E CCC P CAG CCT Q P 2431 AAG GAG K E 2491 AGA AAC (AGA R AGC S GCC A CTT L CTC CCA P GAG E AAG K TTG L CCT P CAG Q AAC N CAG Q GAA E AGA R CCT P ACT GCC A CAG Q AAG K AAA K CCG P GCC TTG CCA P AGA R CCC P GAG E GCC A ပ္ပိပ္သ CCT CTG L CAT H ACA T TCT TCC S CTC AGG R CTC L ACA T AAG K TCC S CCG 2281 CAG ACC Q T 2341 GAG GAG E E E CCA AAC P N 2461 CCT CTG P L

FIG. 6D

TIC GTG V CAG Q AAT GAC ACC X X X CTG L AAG CAT GGC CAG K H G Q TAA CCC TGC CCT. CCA CCG CCG GCT CCA CAC TGC CGG AAA GCA 3121 CTG CTC GGT GCA CGA TGC TGC CCT GMA AAC ACA GGC TCA GCC GTT CCC AGG GGA TTG AAG CAG CTG AAA AAA CAT CCG CCA GCA NGG GGA GAG CAG CTT CCA GCC TGG G'IC AGA AGG GGT GGG CGA GCC CTT CGG CCC CTC GTC V TTG GAC D TTG TCT 00 K 3211 TCG CAG CAG CAA GGT GGG GGC GAA ACA AAA CAG GGA CAT TTG CAC H GAC CGG GAG TGG AGG GCA A GAG GTG CCA GAC E V P D 3391 GCT TGG ATC AGT GCT A CCC ACC CTG GCC TGG AGC A W S CCT GGG 3331 GAC ATG C CAG 3181 GCC CCC CGG CTC ACA GTG GGA ACC AGG GCC TTA GCC TCC CAG GAT TTC ACA CCT GAG CCC TGC GCC CCT GCT CCA GTC GAG TTG GCC TGA CCC CAC H AGC 5 GAA GAG ACA GAA GGA GGA TGG CAG GAG AGC S TAC GAA E AGC S GGT G ပ္ပိပ္ပ AAT N GAT D GTT V ပ္ပ်ပ္ပ် ပ CCT P CCT P CTT ACT GAG AAC AGG CCC E N R P ეეე ე 2941 AAA GAC | K D | 1001 TGC ACA (

Fig. 6E

3481 CTC CAG GCT GCT GTG AGA GTG TCA AGT GTG TAA GGG CCC AAA CTC AGG TTC AGT GCA GAA 3631 TAG GGG AAA AGA ATG CCT GAC CCT GGG AAG TGC AAC ACA GGT CCT GAG TTG ACC CCC CAG ACA 3811 CCC CAG AGC AGG CCC CGA GCC CCT TCA 3931 GTG GAT GGA CAT GAT CAG TGC TAA GGA AAG TCT GTA AAG GTG AAG GGT TIT GCA CIC IGA IGA CCT CAA AGC ACT TIC AIG GCT GCC CIC IGG CAG TCC AGC GTG CTG GTT CCC CTG GCC AGA GGG TAC TTG CCT 3571 CCA GGT CAG CAG GTA TGC CCG CCC GTA GGT TAA GGG GGC CCT CTA AAC CCC GAA GTG GTT CTC AGT CAC CCT GCC CTG CCC CCC TCA CCA CTA TCA GTG 3751 GTA AGT 3871 TCA GTT (3691 TAG AAT ACA CCA CAC TTT TCA GGT TGT CCA GCT CAC CCC TTT TGG GTG AAG GAC CAA AGA AGG TGT CTT ပ္ပင္ပ GGC CCA GCA CTG CCC CAG ACT CGC TGG CAC GCT CAG CAT CAC ACT GAC ACT GAT GCA GGA TAT GCC TGA CAG GAA CAG TCT TTG CTG CTG GCT ACC ACT CAG CAG AGA GAG ACG TCC GGC GCC CCA ပ္ပပ္ပ TGC CGA CGG CAC CTC ACC TGG GCT CCC TGG TCT GGT TCA TGT GCC CCT CAG AGC ACA 3601

FIG. 6F

GGC AGG	GCA	ອອອ	CÀG	TGA	CAC	ŢĠŢ	AGG	AGC	41/1 GCA GGG CAG TGA CAC TGT AGG AGC ATA GCA AGC CAG GAG ATG GGG TGA AGG GAC	GCA	AGC	CAG	GAG	ATG	999	TGA	AGG	GAC	
4201 ACA GTC		AGC	AGC TGT	CCA	CCA CAT	GCA	TGT	GAC	4231 TCC TCA AAC CTC TTC CAG ATT TCT CTA AGA	ICA	AAC	CTC	TTC	CAG	ATT	TCT	CTA	AGA	
4261 ATA GCA	200	CCC CCT TCC CCA TTG	TCC	CCA	TTG	ນນ	CCC CAG	CTT	4291 AGC CIC TIC TCC CAG GGG AGC TAC TCA	CIC	TTC	TCC	CAG	999	AGC	TAC	TCA	GGA	
4321 CTC ACG	TAG	TAG CAT	TAA	ATC	TAA ATC AGC TGT GAA TCG	TGT	GAA	TCG	4351 . TCA GGG GGT GTC TGC TAG	ງຕູ	GGT	GTC	TGC	TAG	CCT CAA	CSA	CCT	CCT	
4381 GGG GCA		GGG GAC GCC GAG ACT	သည	GAG	ACT	ນວວ	TGG	GAG	4411 AAG CTC ATT CCC ACA TCT TGC CAA GAC	OTC .	ATT	ນູ	ACA	TCT	TGC	CAA		AGC	
4441 CTT TGT		CCA GCT GTC CAC ATT GAG	GTC	CAC	ATT	GAG	TCA	GAC	4471 TGC TCC	ງບູ	່ງງວ	GGA	GGA GAG AGC CCC GGC	AGC	ວິນ	ວຍູຍ	CCC	CAG	
4501 CAC ATA	AAG	AAC	TGC	AGC	AAC TGC AGC CTT GGT ACT GCA	GGT	ACT	GCA	4531 GAG TCT	ict (່ງຄວ	rTG	TIG TAG AGA ACT CTT	AGA 2	ACT (CTT	TGT AAG	AAG	
4561 CAA TAA AGT ITG GGG TGA TGA CAA ATG TTA AAA	AGT	TTG	999	TGA	TGA	CAA	ATG	TTA	AAA A	AAA 45	4596								

FIG. 6G

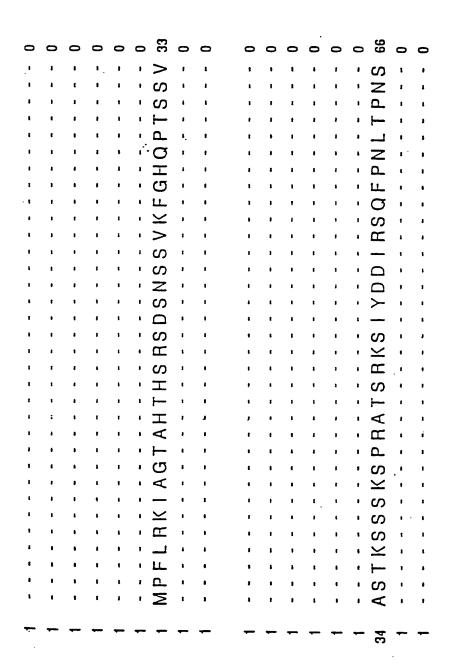


FIG. 7A

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FIG. 7B



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FIG. 7D

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FIG. 7E

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FIG. 7F

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FIG. 7G

FIG. 7H

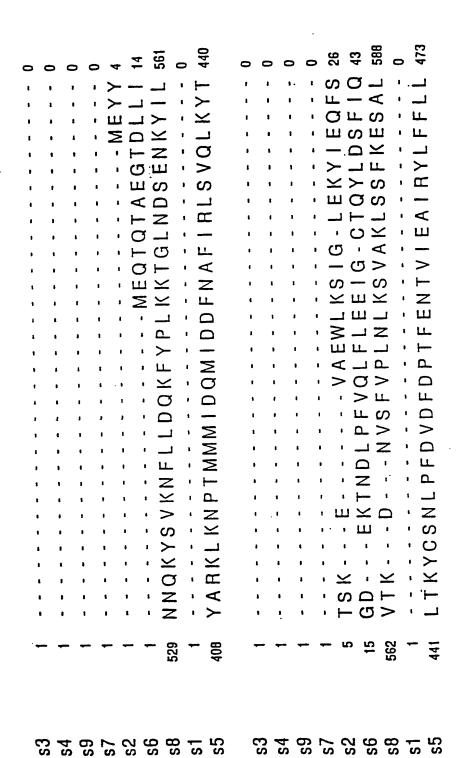


FIG. 7I

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FIG. 7J

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FIG. 7K

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FIG. 7L

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s9	47	AGPAAI
s7	-	
s2	180	KLSS VLPTSTQKRSVRSNNAKPFESY
se	241	LA QNKG I SHNN A EGKL R I DNT E
s8	813	NTSPORTLSTSKONKPIRLVRASTKISRSKRS
s 1	112	TTTEQPKPAVQTKGRPHSQCLNSSPLSHAQLM
s5	203	TDPFDQHLD - EKNNEVFELEVALSSLGALVVL

FIG. 7M

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s2	207	SELINSRISD
se	269	RIBOIFN ORPPSEFISTNLAG -
s8	846	PPOLLSSPIEASSSSPDSLTSSYTPASTHV
s1	145	PAPSAPCSSAPSVPDISKHRPQA-FVPCKIPS
s5	735	PGEP V VWDGP V Y K L PGNN L F AS N EM
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FIG. 7N

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	HRPKPSRPVPGSTSKLGDA
	KRCSLNSIESSLQKINKAYYKLTYTVLNNYKG
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	MALLQGLDQTRLSSKL
	RVVSSTSAADIFEENDITFADAPPMFDSDD
	KSSMTLDLGS AS RCDDS FGGGGNSGNAVIPSD.
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FIG. 70

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FIG. 7P

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FIG. 70

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FIG. 7R

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FIG. 7U

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FIG. 7V

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FIG. 7W

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FIG. 7X



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FIG. 7Y

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FIG. 7Z



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FIG. 7AA



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